

-continued

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tttttttggg acttttgcta gatataat tttt acacatactc atttttatga gtcttaagt	360
caatacgttg gtaacggaat actgggtatt tgctattcct tccttgctgt acctaggttg	420
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agaaataaga gagtaatcag gagagagaaa gagagaaaag attgagtaat ctgtaagaca	540
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acagggtatta gttcaatcca gataaaaaat gaagtgttaa aagacataga agaaaaactt	660
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1-45. (canceled)

46. A method for controlling corn rootworm infestation on a plant, comprising topically applying to the plant a pesticide composition comprising a dsRNA targeting for suppression an essential gene in the corn rootworm and providing the plant in the diet of the corn rootworm, wherein the plant comprises a nucleic acid sequence encoding at least a first *Bacillus thuringiensis* insecticidal protein for controlling the corn rootworm.

47. The method of claim 46, wherein the *Bacillus thuringiensis* insecticidal protein, or proteins, is selected from the group consisting of a Cry3, a TIC851, a CryET70, a Cry22, a binary insecticidal protein CryET33 and CryET34, a binary insecticidal protein CryET80 and CryET76, a binary insecticidal protein TIC100 and TIC101, a binary insecticidal protein PS149B1, a VIP insecticidal protein, a TIC900 or related protein, a TIC901, a TIC1201, a TIC407, a TIC417, and an insecticidal chimera thereof.

48. The method of claim 46, wherein the dsRNA is topically applied using a spray mixer, by expressing the dsRNA in a microbe and applying the microbe onto the plant, or by applying the dsRNA in a topical composition.

49. The method of claim 48, wherein the topical composition further comprises a *Bacillus thuringiensis* insecticidal protein.

50. The method of claim 46, wherein the composition comprises dsRNA encapsulated in a synthetic matrix and applied to the plant surface or a seed coating.

51. The method of claim 50, wherein said matrix is a polymer.

52. The method of claim 50, wherein the composition is topically applied as a microbe that is engineered to express the dsRNA, or comprises a fermentation product from the microbe.

53. The method of claim 46, wherein the dsRNA is transcribed from a DNA sequence of at least about 19 contiguous nucleotides selected from the group consisting of SEQ ID NO:1-143, SEQ ID NO:169-174, and the complement thereof.

54. The method of claim 46, wherein the corn rootworm is selected from the group consisting of *Diabrotica virgifera*, *Diabrotica barberi*, and *Diabrotica undecimpunctata*.

55. The method of claim 46, wherein the diet is selected from the group consisting of a plant cell, a plurality of plant cells, a plant tissue, a plant root, and a plant seed, wherein the diet comprises a pest inhibitory amount of the dsRNA.

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